



45<sup>th</sup> International Conference on  
Environmental Systems

# Miniature Liquid Chromatographic Systems for Human and Robotic Missions

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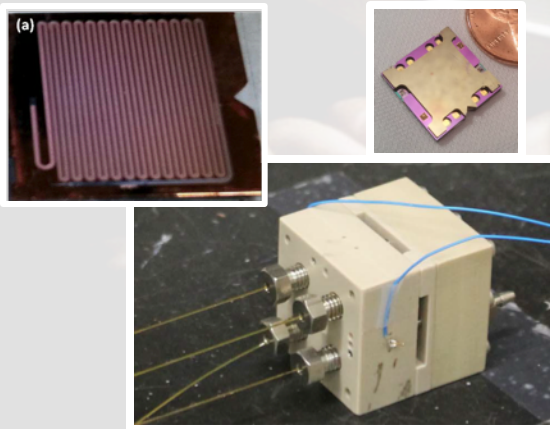
- SMD (Science Mission Directorate): Our instruments are meant to address two fundamental questions in Astrobiology, namely, "How does life begin and evolve?" and "Does life exist elsewhere in the Universe?" For these reasons, the *Science Objectives* for the analyzer are to look for:
  - ✓ Signs of extinct life by detecting Carboxylic Acids and Lipids - the longevity and preservation of carboxylic acids and lipids offer a chemical insight into potential primordial biological activity.
  - ✓ Extant life by searching for Peptides and Proteins - macromolecules that strongly indicate a biotic origin.
- Provide organic molecular detection and life detection capabilities for future landed missions to Mars, Europa, Titan, Enceladus, and other planetary bodies.
- Astrobiology field research on Earth.
- HEOMD (Human Exploration & Operations Mission Directorate): Experiments on ISS, astronaut health monitoring, environmental monitoring.
- Of the three main chromatographic technologies (GC, CE, LC), liquid chromatography is the least advanced.



### Gas Chromatography (GC)

- Can be coupled to a MS
- Very fast
- Can separate volatile small molecules
- Need to derivatize amino acids, fatty acids, etc.

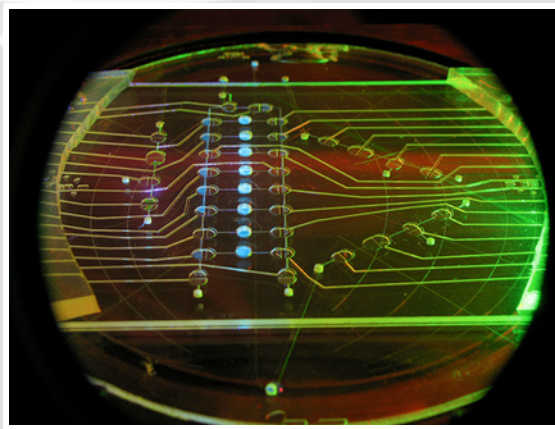
Madzunkov, S.M., MacAskill, J.A., Simcic, J., Kidd, R.D. & Darrach, M. (2013). Recent developments in gas chromatographs and mass spectrometers for crewed and robotic space missions. *J. Am. Inst. Aeronautics & Astronautics*. Electronic 2012-3453.



### Capillary Electrophoresis (CE)

- Coupled to laser-based fluorescence detector
- Very sensitive/specific
- Can separate small molecules to macromolecules

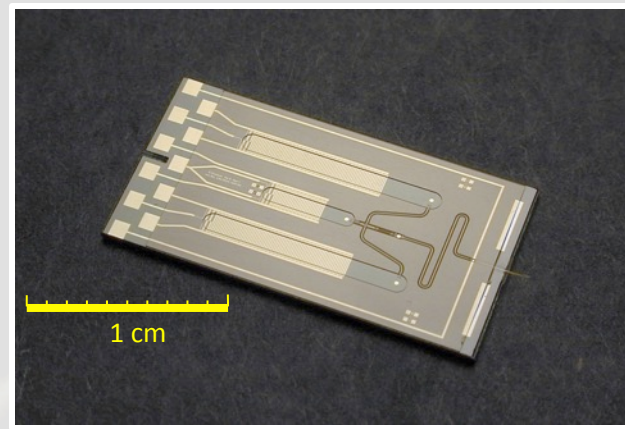
Willis, P. A., Stockton, A. M., Microchip Capillary Electrophoresis for In Situ Planetary Exploration. In *Capillary Electrophoresis and Microchip Capillary Electrophoresis*, John Wiley & Sons, Inc.: **2013**; pp 277-291.



### High Performance Liquid Chromatography (HPLC)

- Can be coupled to a MS
- Well suited for non-volatiles
- Can separate small molecules to macromolecules
- No derivatization required

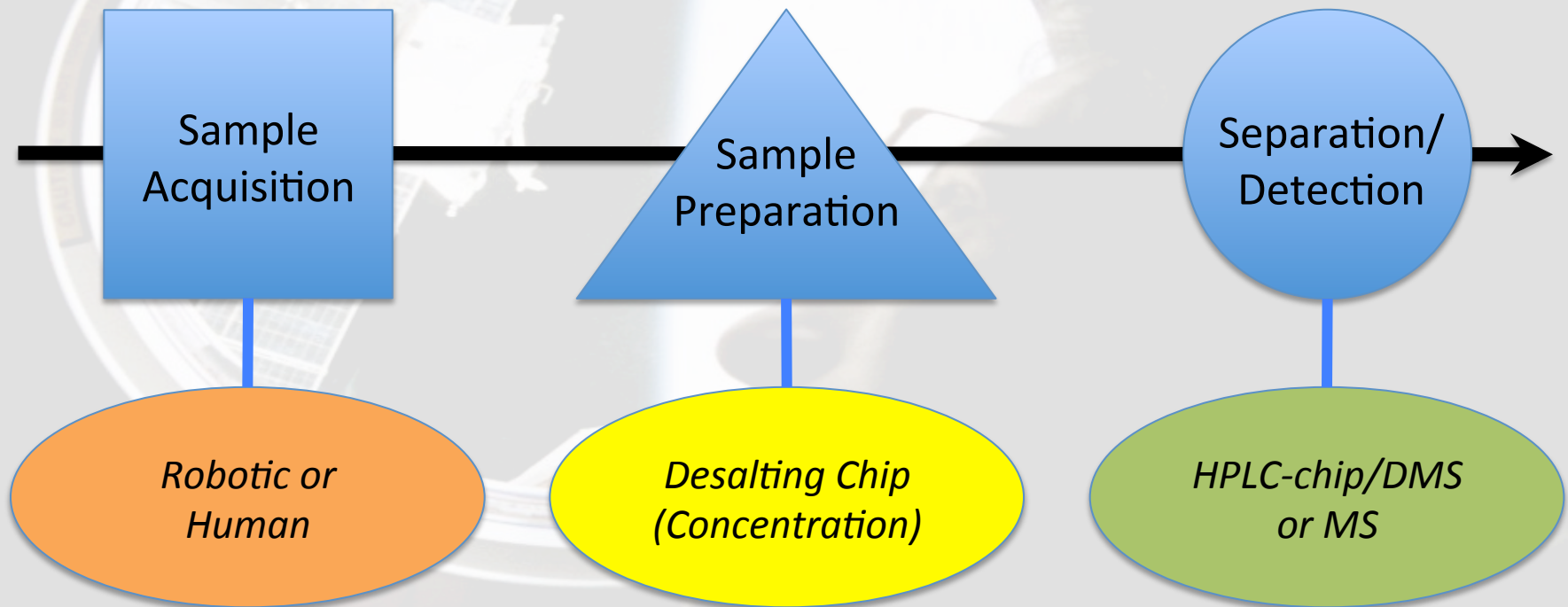
Xie, J., Miao, Y., Shih, J., Tai, Y.-C. & Lee, T.D. (2005). Microfluidic platform for liquid chromatography-tandem mass spectrometry analyses of complex peptide mixtures. *Anal. Chem.* **77**, 6947-6953.







In designing *in situ*, liquid-based analytical instruments, three areas need to be addressed: *Sample Acquisition*, *Sample Preparation* and *Separation/Detection*.

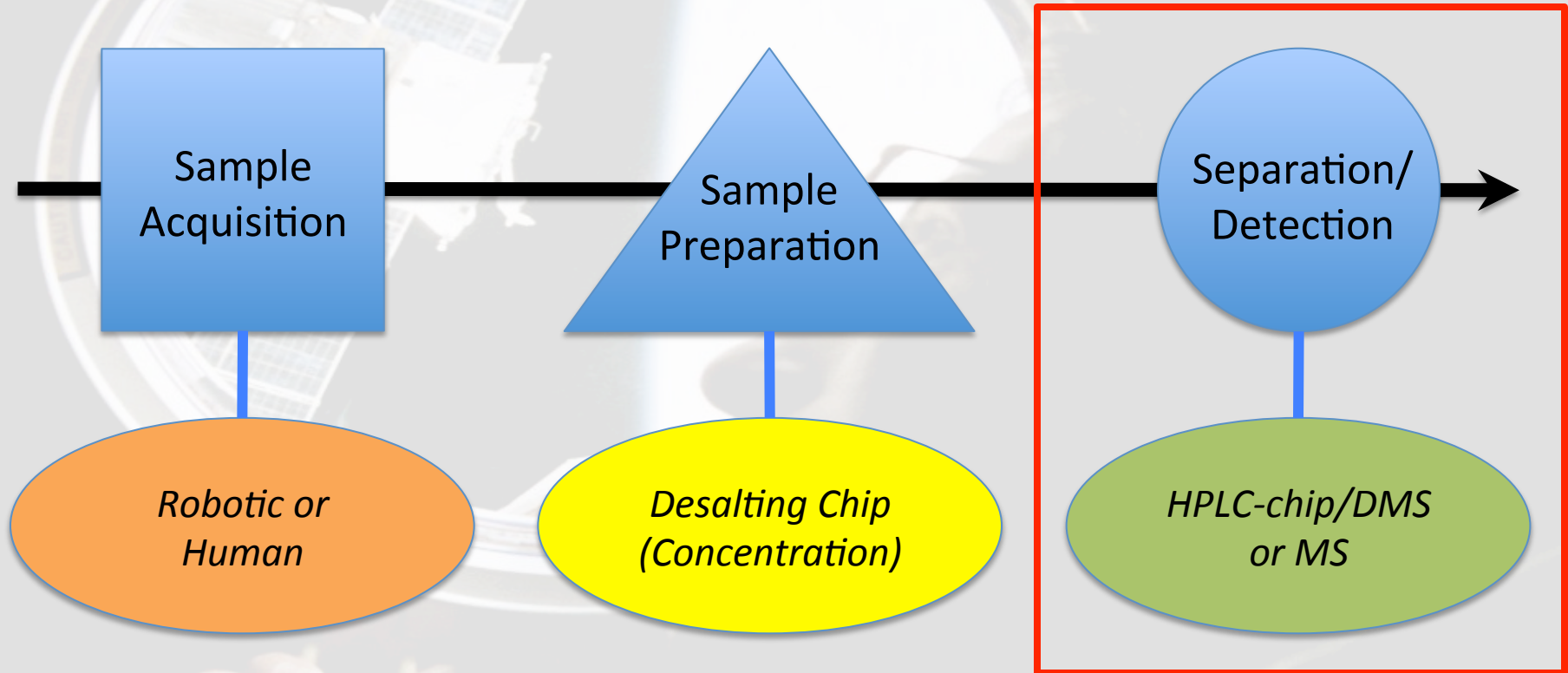


We are pursuing research into all three areas focusing, thus far, on developing a unique, miniaturized solute analyzer based on microfluidics technology. This analyzer consists of an integrated microfluidics High Performance Liquid Chromatographic-chips coupled to either a Differential Mobility Spectrometer ( $\mu$ HPLC-chip/DMS) or Paul Ion Trap Mass Spectrometer ( $\mu$ HPLC-chip/MS)





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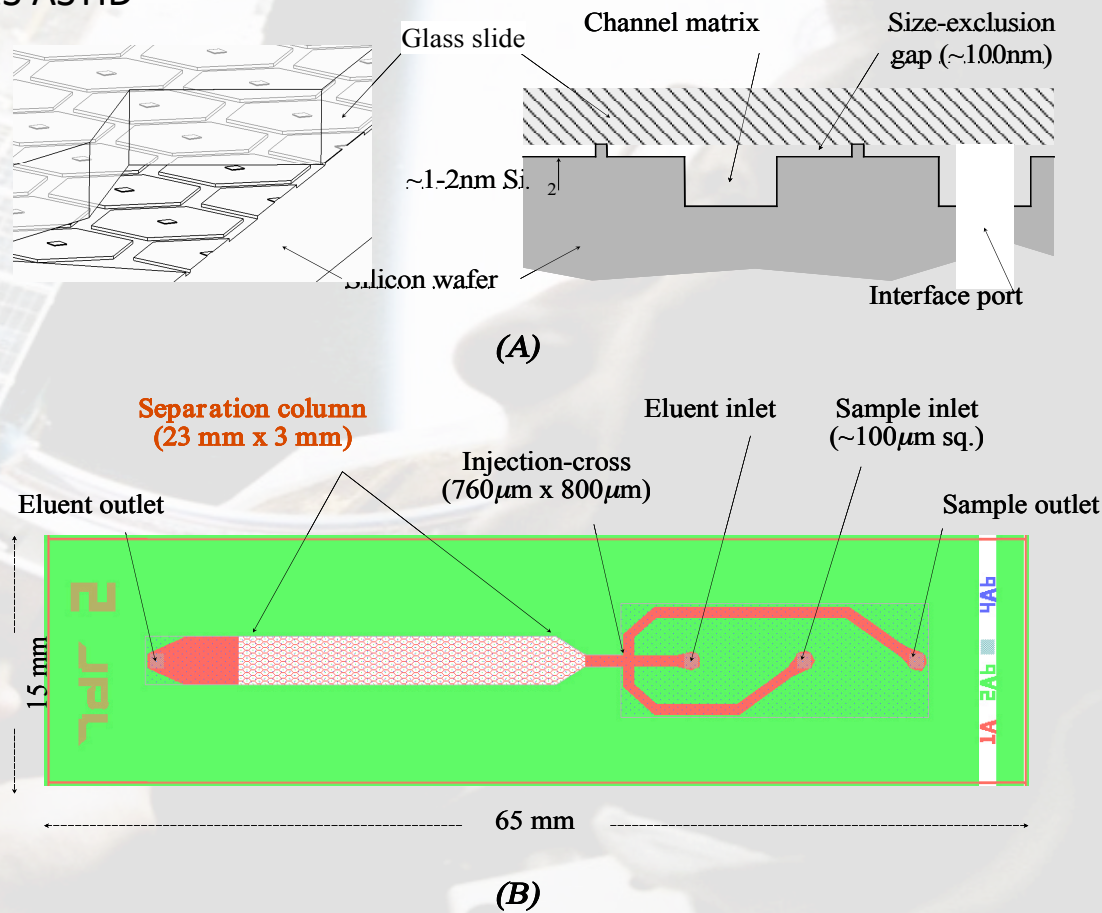


## *nanoSEC-chip/Laser-Induced Fluorescence Detector*

Collaborators: JPL's Micro Devices Laboratory

Target: fatty acids, small peptides, carbohydrates, kerogenic and humic fractions

Funding: NASA ROSES ASTID

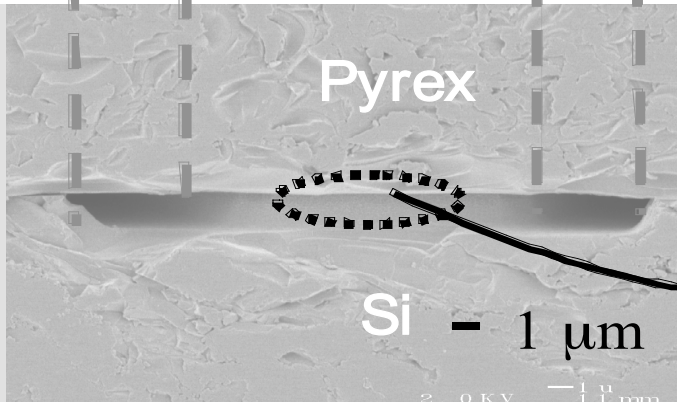
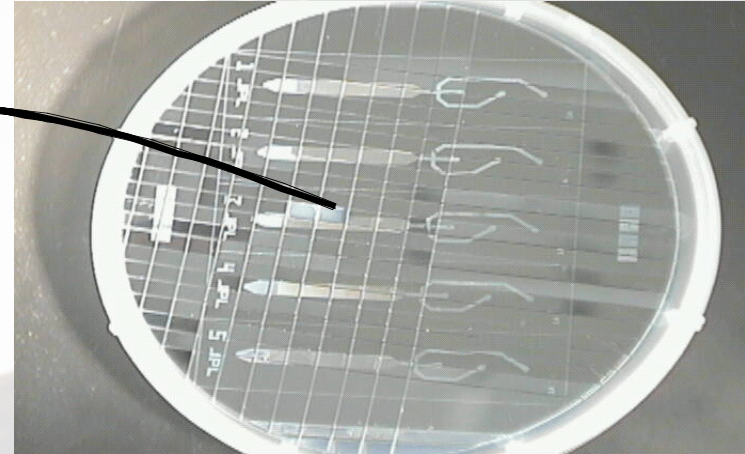
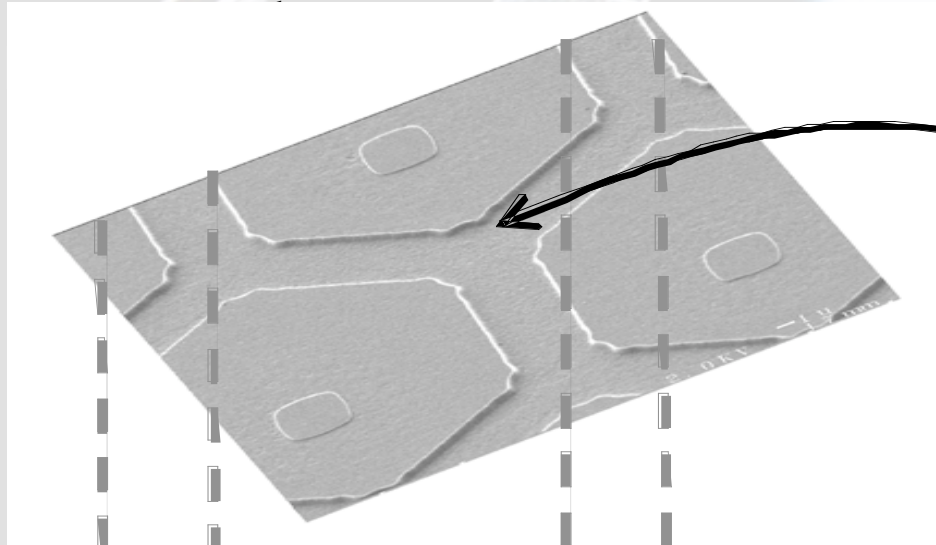


**Figure 1: (A) 3-D and cross-sectional views of nSEC column schematics,  
(B) Top view of nSEC device layout**

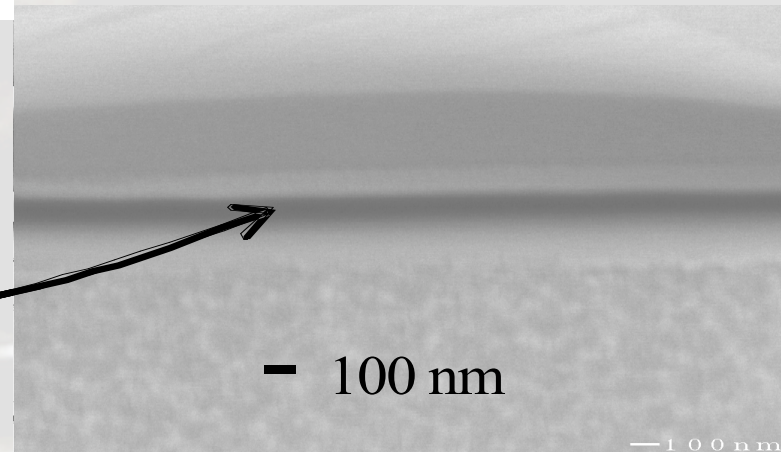


Top view of nSEC channels

Sealed Si wafer containing five nSEC



Cross-section of bonded wafer  
showing 1  $\mu\text{m}$  and 100 nm gaps



SEM image confirming presence of  
100 nm gap in sealed nSEC device



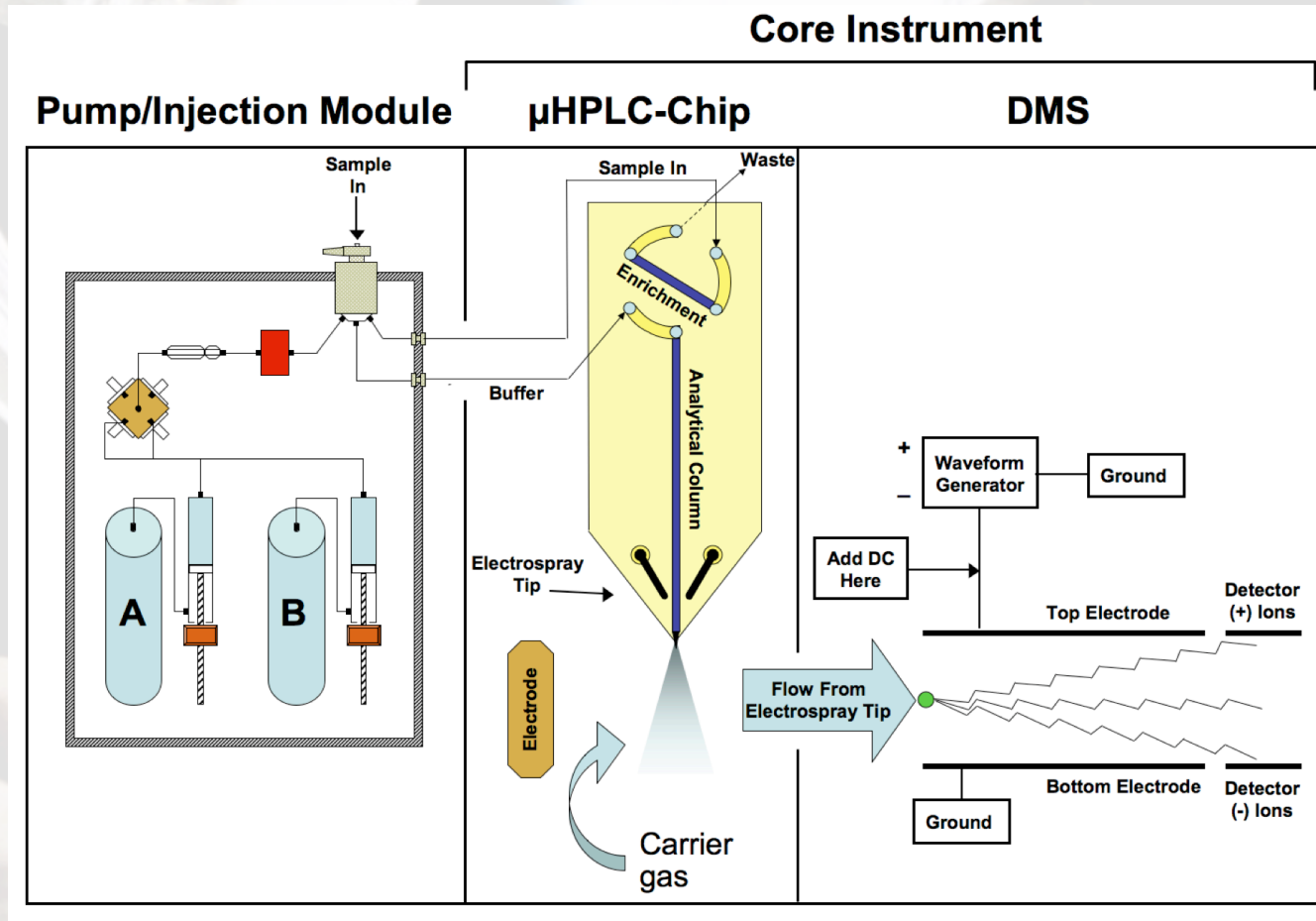


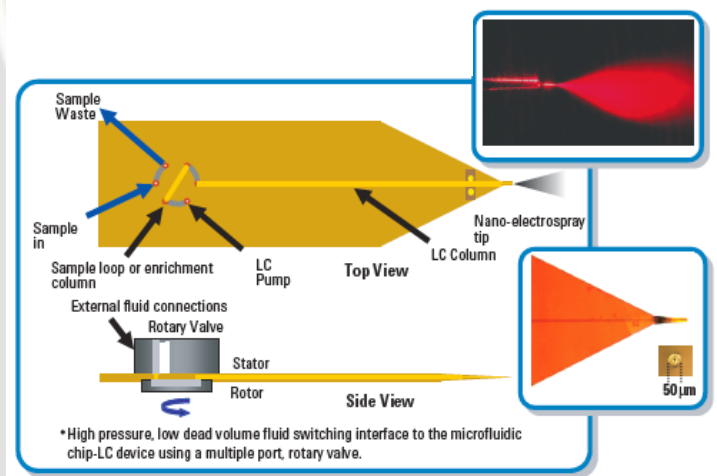
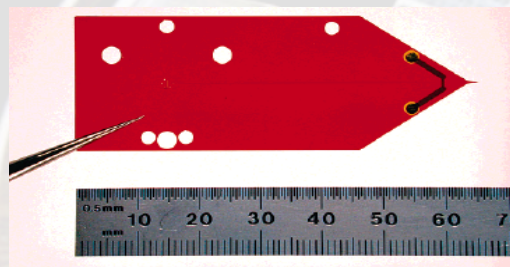
## *HPLC-chip/Differential Mobility Spectrometer (DMS)*

Collaborators: Agilent's Molecular Separations Lab & Sionex LLC

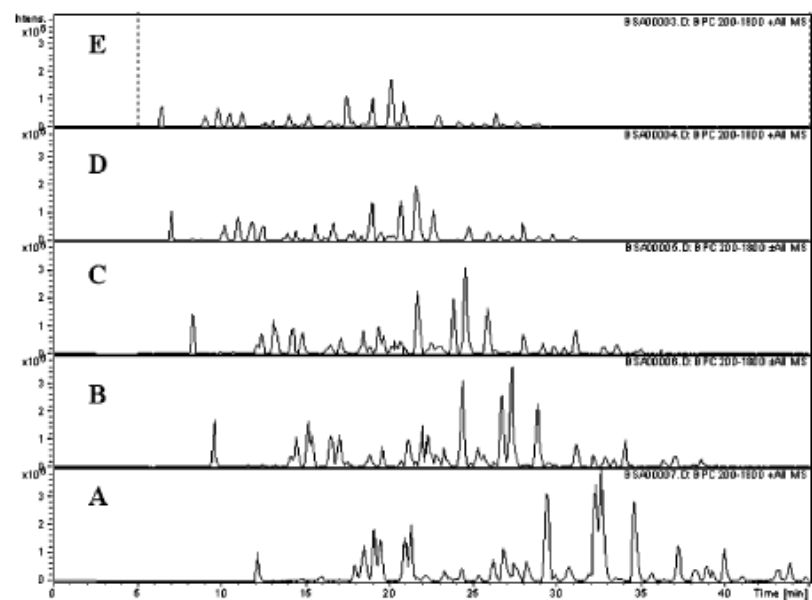
Target: fatty acids, small peptides, carbohydrates

Funding: NASA ROSES PIDDP

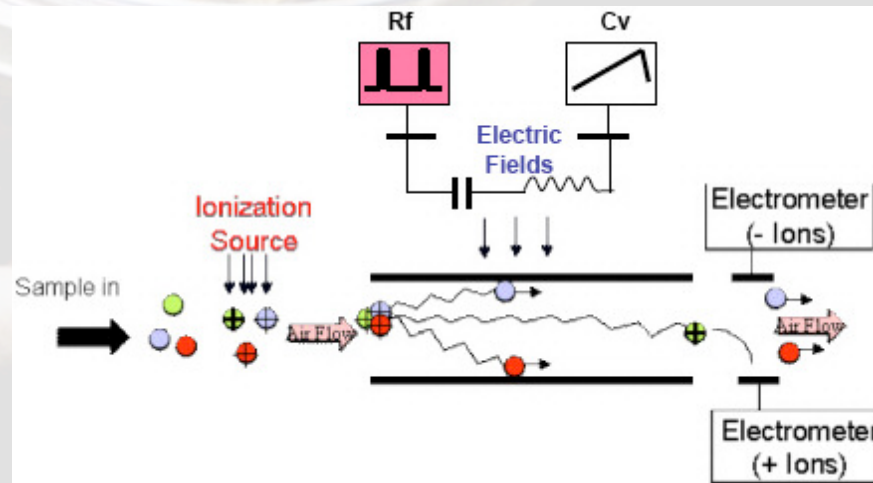
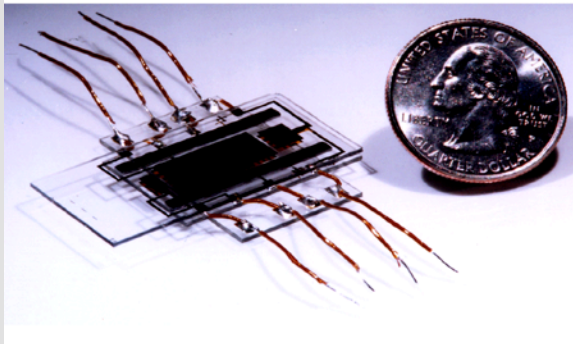
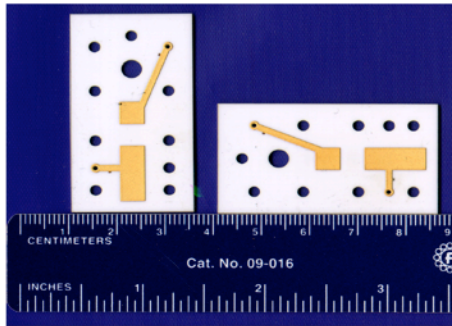




(Top) Photograph of the  $\mu$ HPLC-chip. The holes were used for alignment purposes. The dark pattern on the right end of the chip is the electrodeposited metal for contact to the fluid flow channel near the electrospray tip. (Bottom) Schematic top and side views of the  $\mu$ HPLC-chip



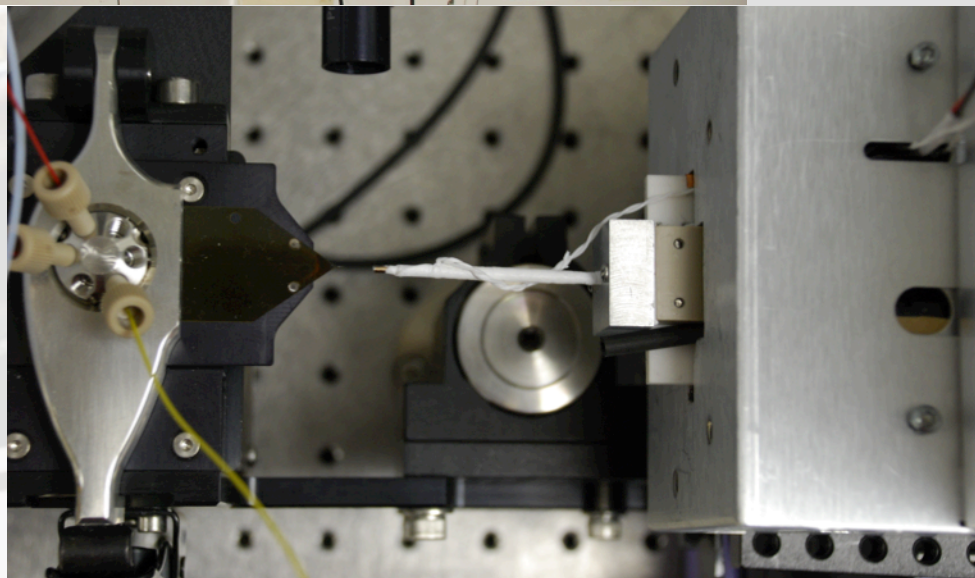
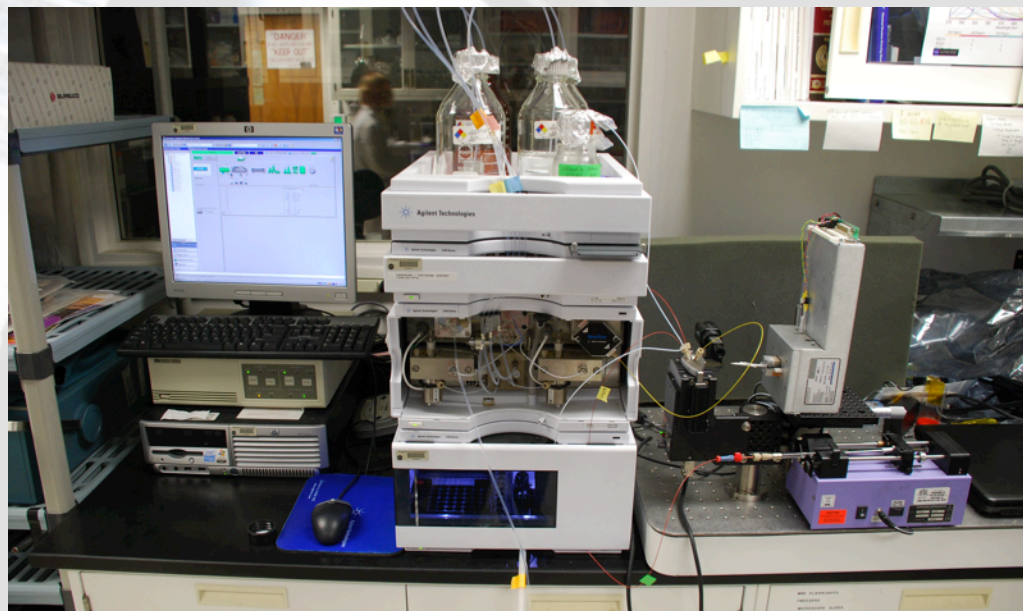
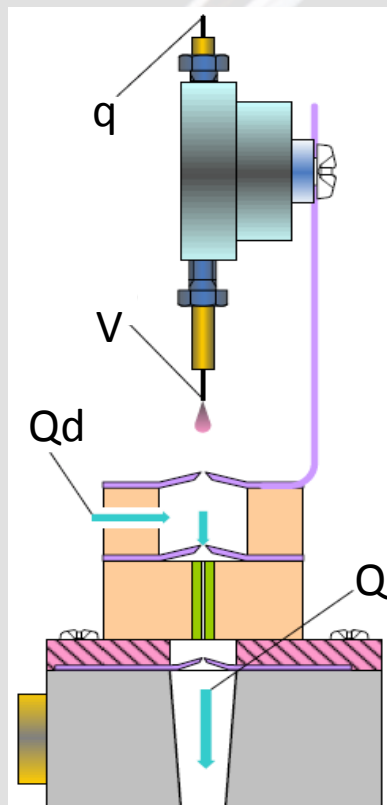
Base peak chromatograms of a 20 fmol tryptic digest of bovine serum albumin (BSA) running under different LC flow rate: (A) 100, (B) 150, (C) 200, (D) 300, and (E) 400 nL/min. The vertical scales in each spectrum are identical.







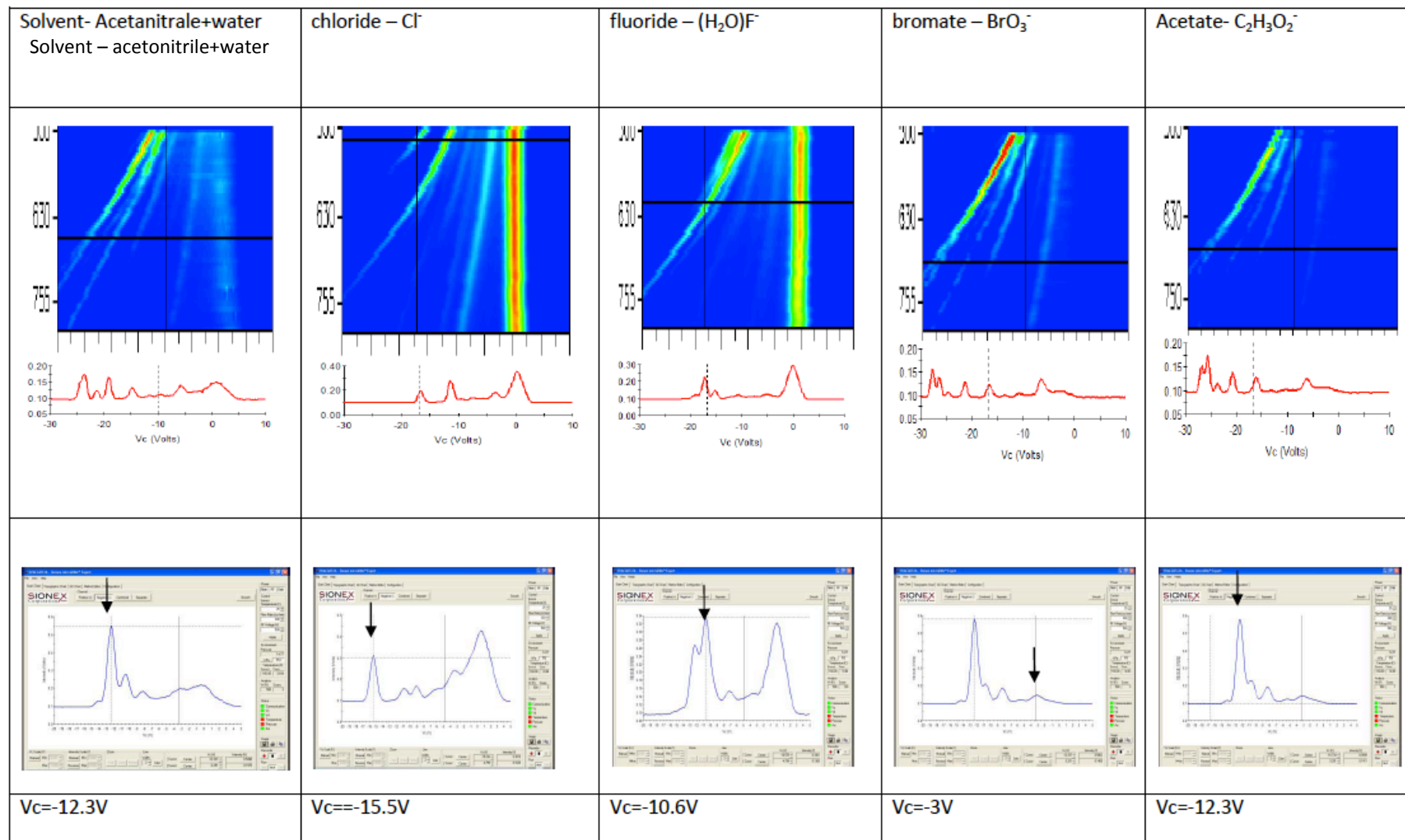
## Preliminary HPLC electrospray to DMS



Coy, S.L. Krylov, E.V., Nazarov, E.G., Fornace Jr., A.J. & Kidd, R.D. (2013). Differential mobility spectrometry with nanospray ion source as a compact detector for small organics and inorganics. *Int. J. Ion Mobil. Spec.* 16, 217-227.



Target ions: fluoride – 37/19; chloride – 35; formate – 45; acetate – 59; bromate – 127/129. Mixture – 50+10+10+10+10 ppm. Q = 1 L/min; Qd = 10 L/hr; V = 1.9 kV; q = 800 nL/min; C<sub>V</sub> = -25...5 V.





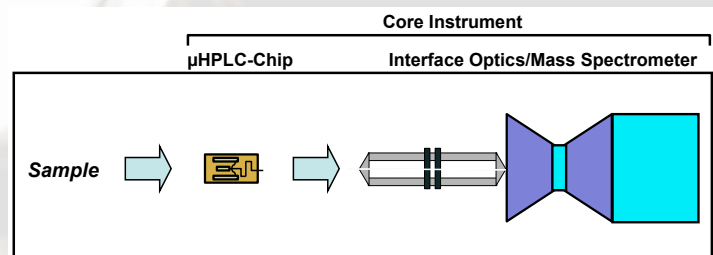
## HPLC-chip/Mass Spectrometer (MS)

Co-Investigator: Y.-C. Tai, Caltech

Target: Carboxylic Acids & Lipids (fatty acids and sterols), Amino Acids & Peptides

Funding: NASA ROSES ASTID

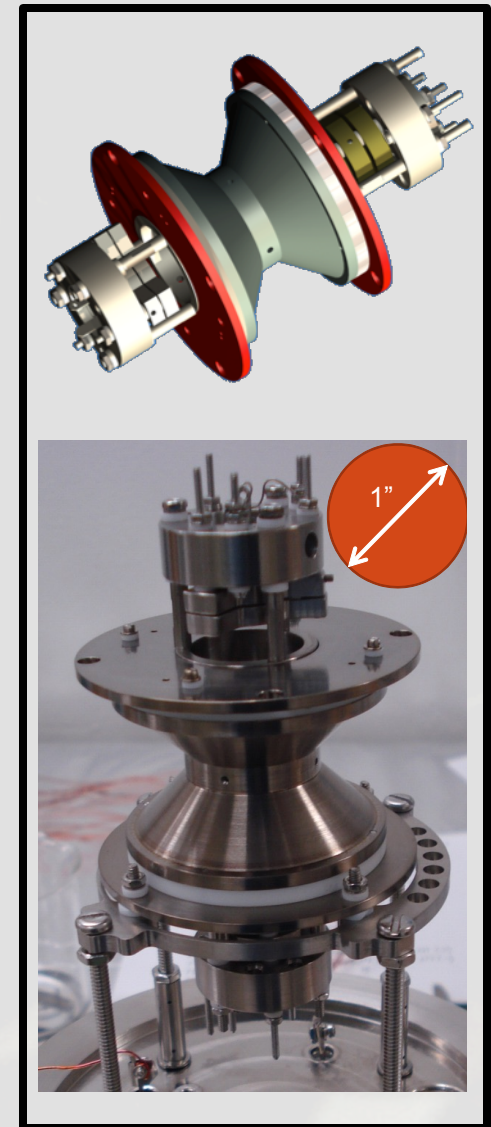
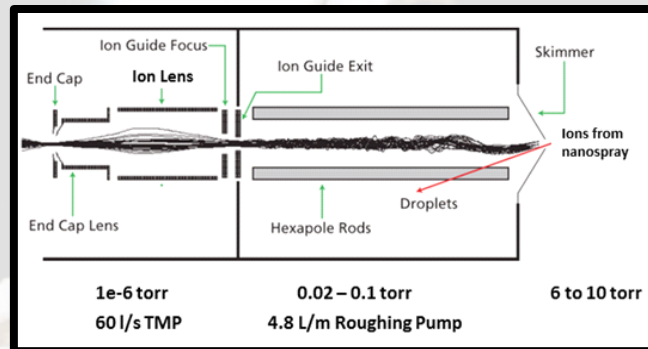
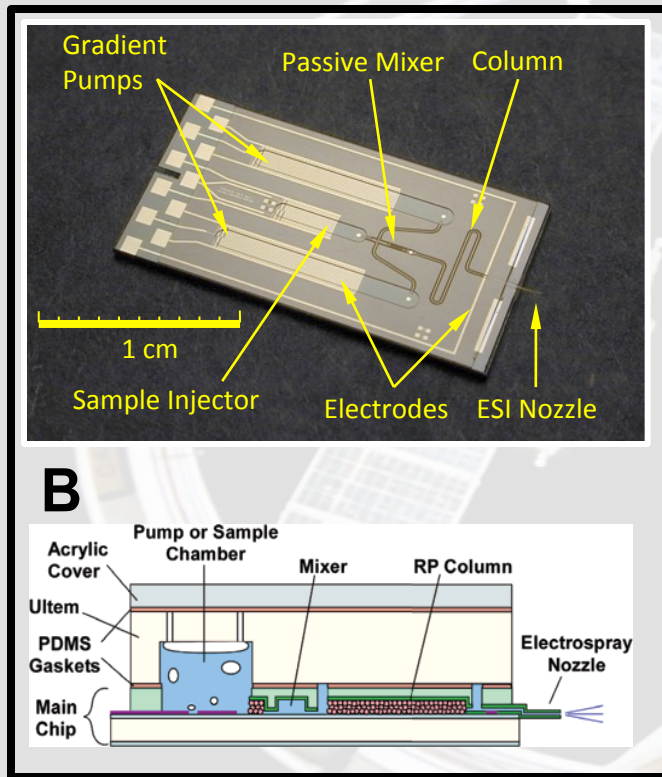
[The ASTID NRA](#) specifically solicits “lab-in-a-teacup” development projects. The goal is to apply micro/nanotechnology to planetary instrumentation and develop highly integrated miniature instrument suites with the capability to address astrobiology questions in planetary exploration.



### Capability Goals:

- **Sensitivity:** Low parts-per-billion/trillion (ppb/ppt) depending on species.
- **Salt Compatibility:** Desalting carried out by a second, ion-exchange HPLC-chip.
- **Resolution:** Capable of separating isomers and determining fatty acid chain lengths.
- **Mass Range:** From small organic acids (e.g. acetate) to macromolecules.
- **Detect Unknowns:** Yes, with no *a priori* selection of chemical species.
- **Mass / Power:** <6 kg / <20 watts for complete instrument based on current parameters for miniaturized HPLC-chip and MS instruments (under our funded PIDDP award).
- **Chemical Derivatization / Labeling:** None required.
- **Expandability:** Several chromatographic chips could be integrated together (see **Figure 1-7**) and serviced by single detection system.





Xie, J., Miao, Y., Shih, J., Tai, Y.-C. & Lee, T.D. (2005). Microfluidic platform for liquid chromatography-tandem mass spectrometry analyses of complex peptide mixtures. *Anal. Chem.* 77, 6947-6953.

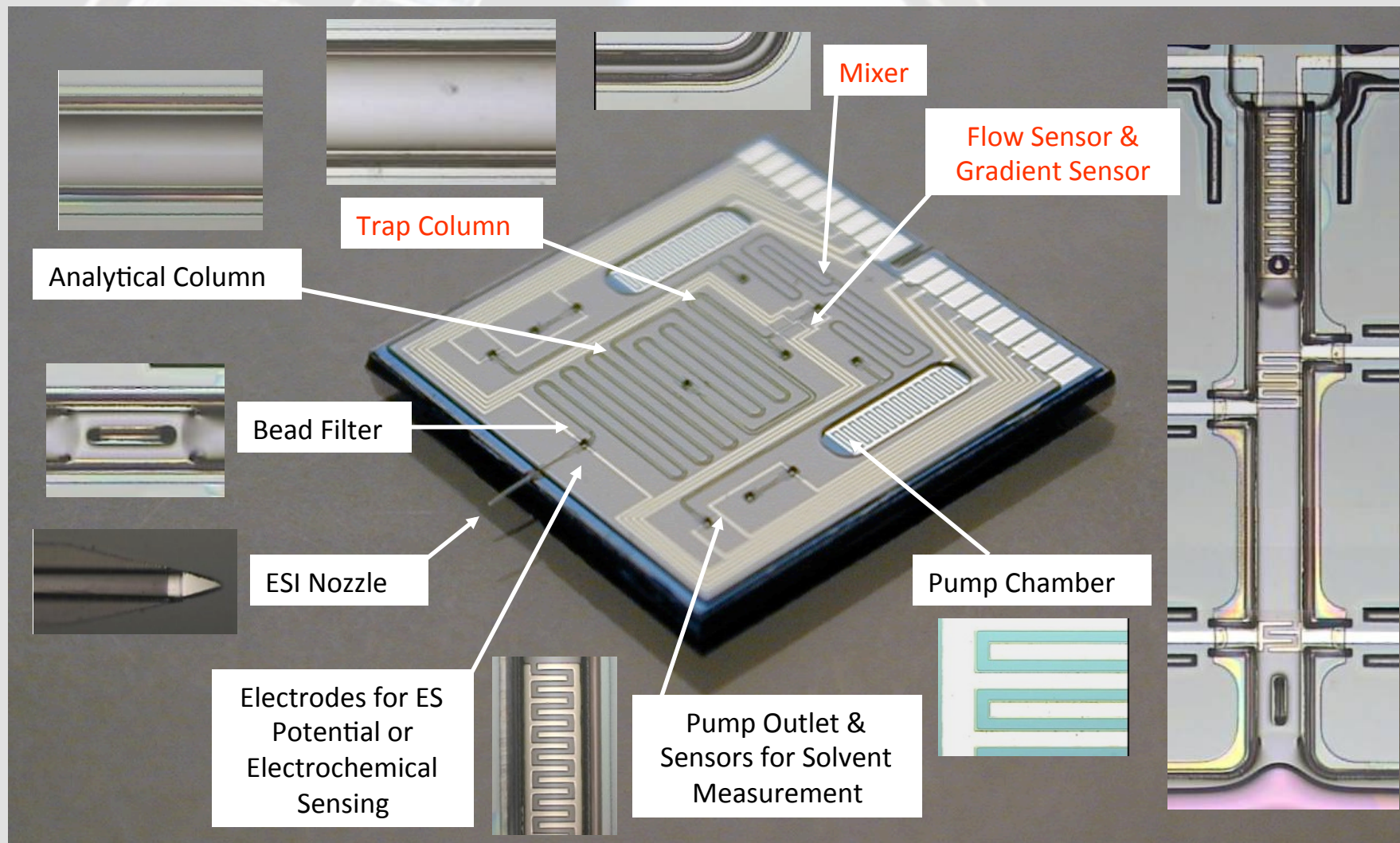
Madzunkov, S.M., MacAskill, J.A., Simcic, J., Kidd, R.D. & Darrach, M. (2013). Recent developments in gas chromatographs and mass spectrometers for crewed and robotic space missions. *J. Am. Inst. Aeronautics & Astronautics*. Electronic 2012-3453.

The figure displays two plots related to the analysis of a protein sample. The top plot is a mass spectrum (MS<sup>2</sup> 653) showing relative intensity (0 to 100) versus mass-to-charge ratio (m/z, 300 to 1100). The spectrum features several prominent peaks labeled b<sub>3</sub><sup>+</sup>, b<sub>4</sub>, y<sub>4</sub>, y<sub>6</sub>, y<sub>7</sub>, y<sub>8</sub>, y<sub>9</sub>, and y<sub>10</sub>. The bottom plot is a chromatogram showing intensity (0 to 5 x 10<sup>7</sup>) versus time. The chromatogram is divided into several phases: Column Wash, Column Equilibration, Sample Injection, Sample Wash, and Start Gradient. The signal is low during the wash and equilibration phases, rises sharply during the sample injection phase, and remains high during the sample wash phase. The signal then decreases during the start gradient phase and remains low throughout the rest of the run.

Mass spectrum of Limonene (m/z 35 to 150). The base peak is at m/z 68. The x-axis is labeled 'm/z' and the y-axis is labeled 'Relative Intensity (%)'. The spectrum shows characteristic peaks for Limonene, including m/z 35, 39, 41, 42, 43, 46, 51, 53, 55, 57, 58, 65, 66, 67, 68, 69, 70, 74, 77, 79, 80, 81, 82, 83, 87, 90, 92, 93, 94, 95, 105, 106, 107, 115, 119, 121, 133, 135, and 136.



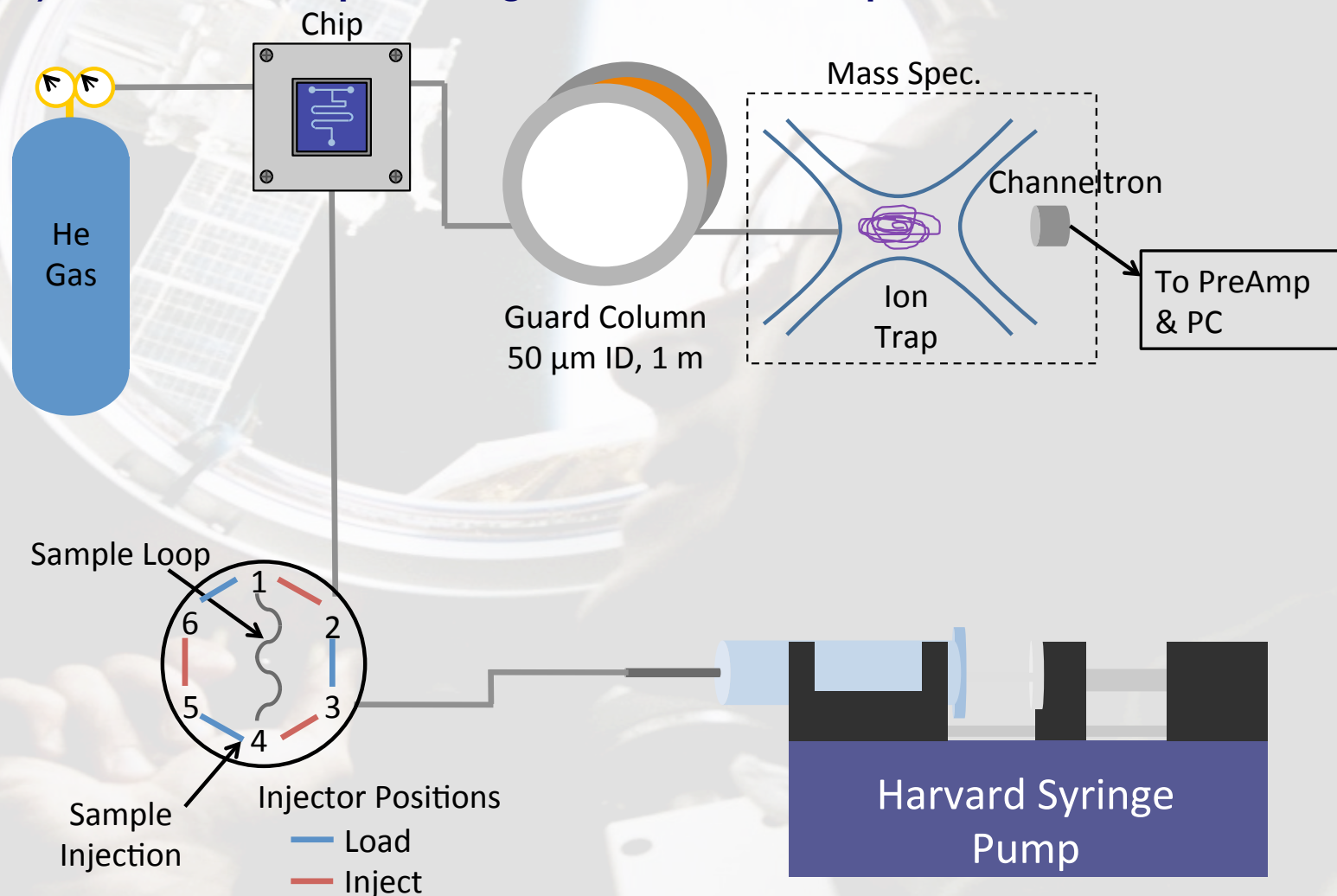
uHPLC





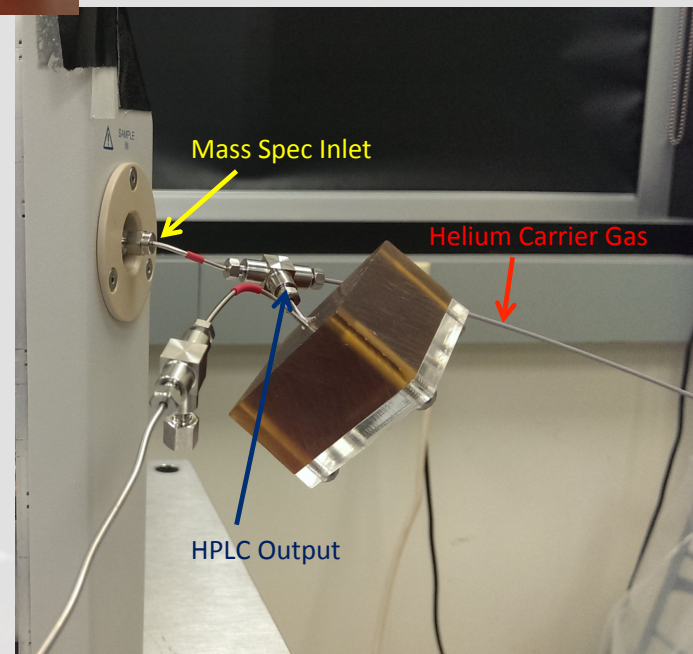
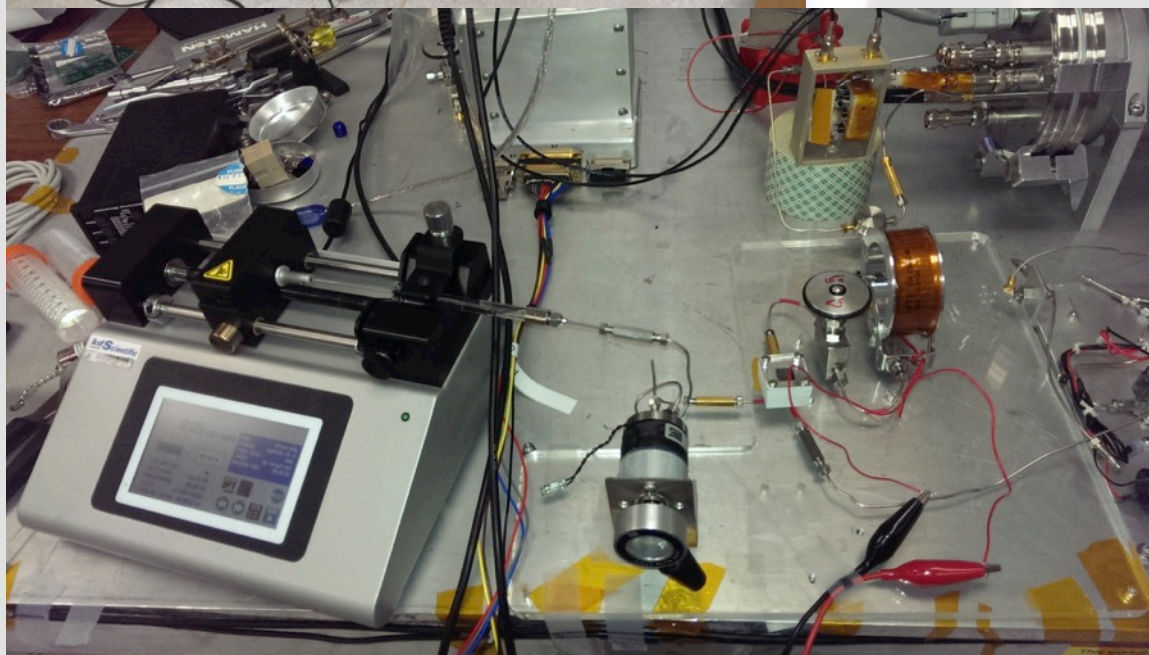
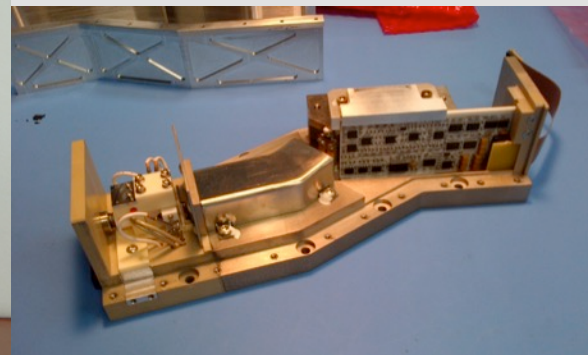
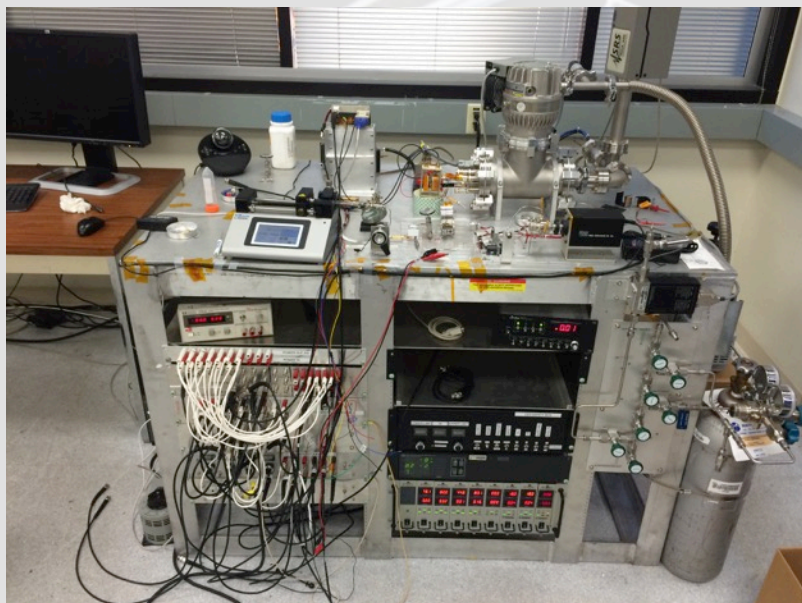


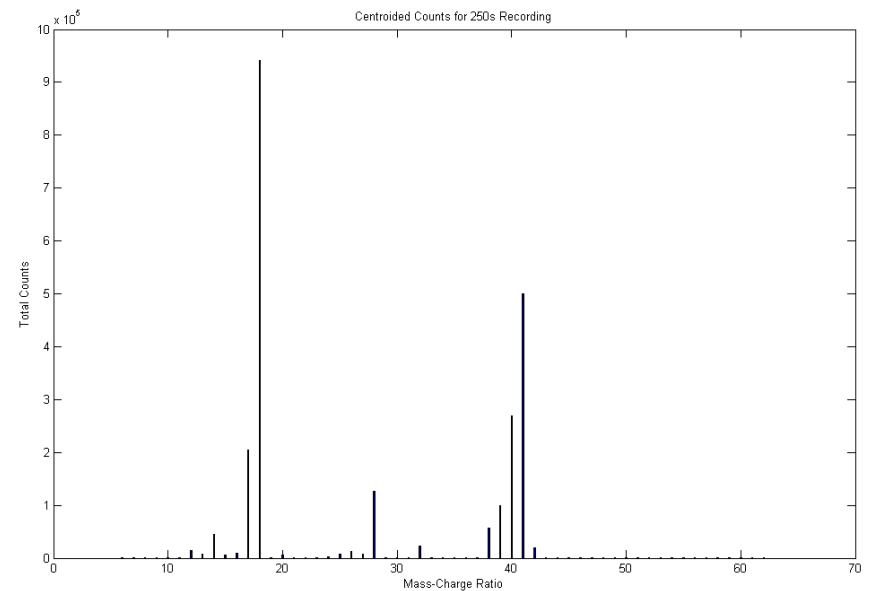
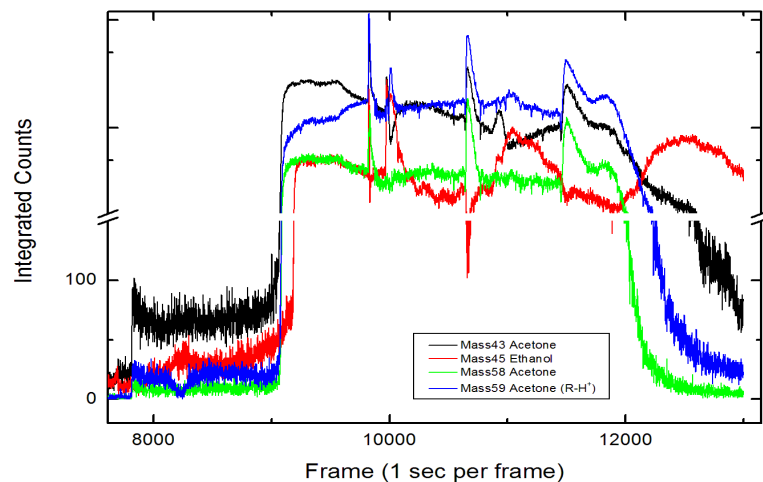
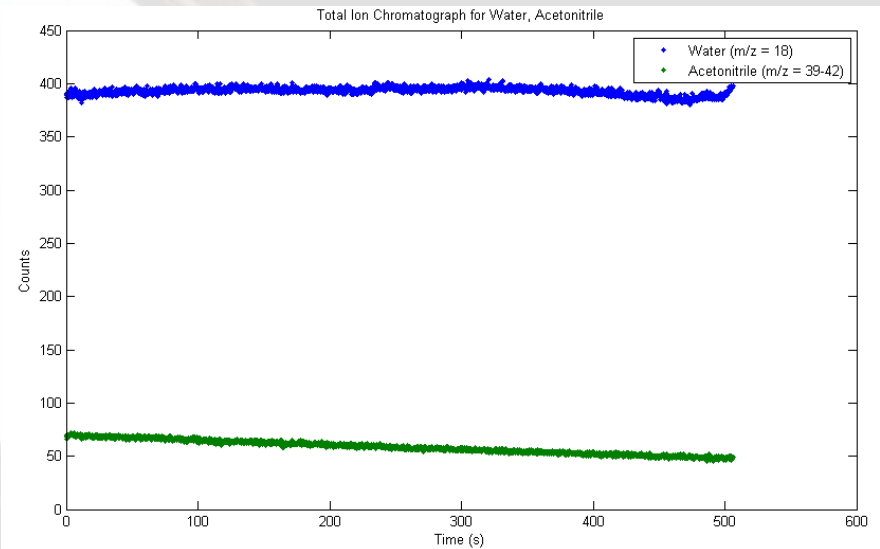
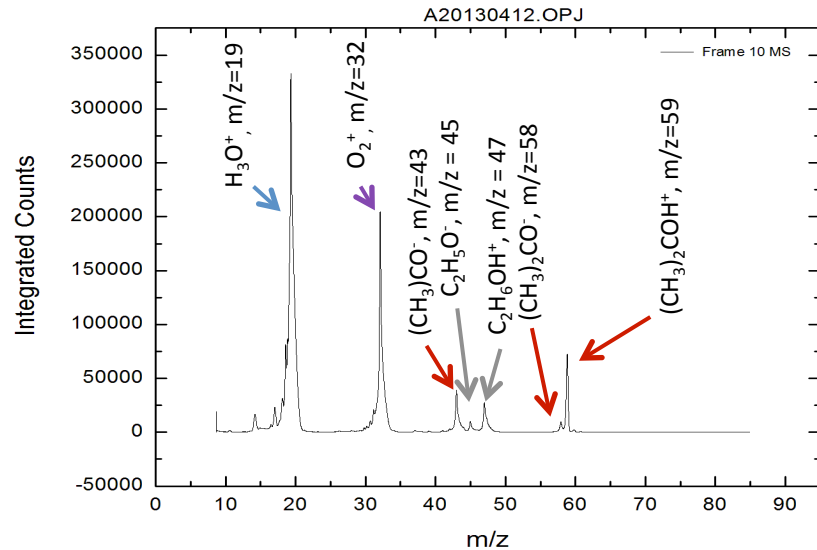
Since the flow rate of the HPLC-chip is so low, ~20-50 nL/min, we explored direct entry (no ESI) into Paul ion trap and magnetic sector mass spectrometers.



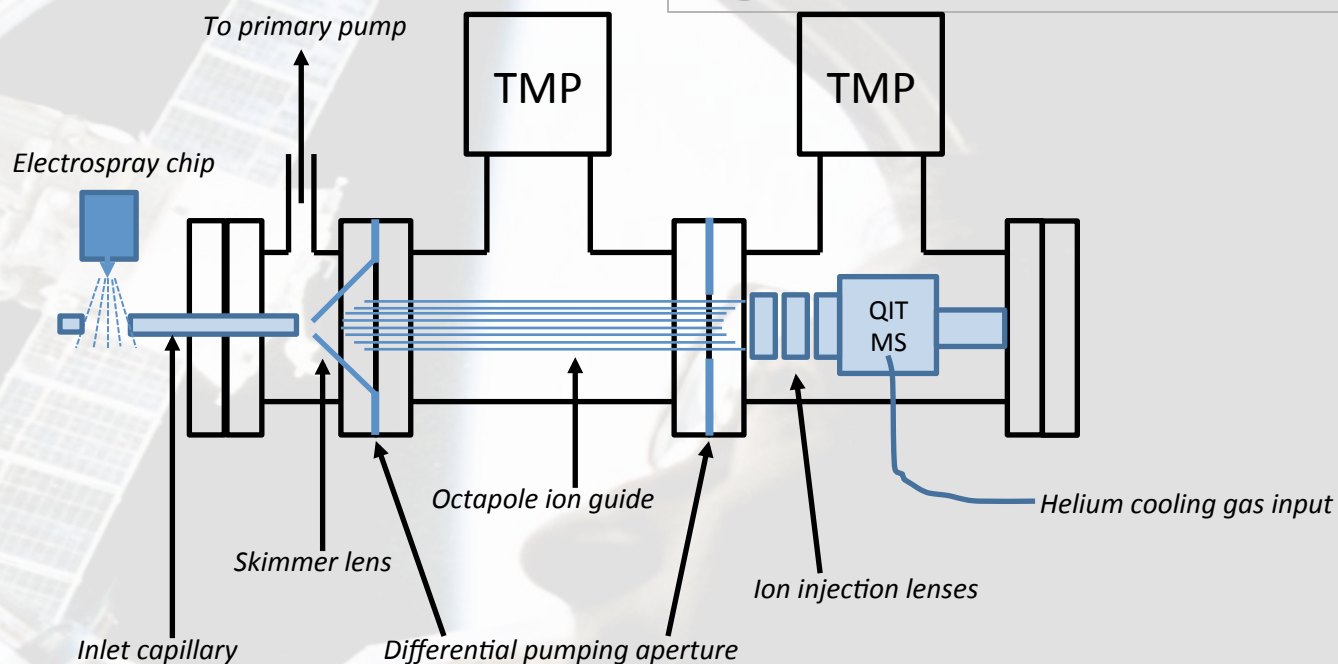


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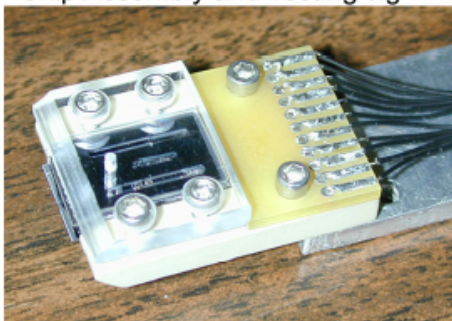






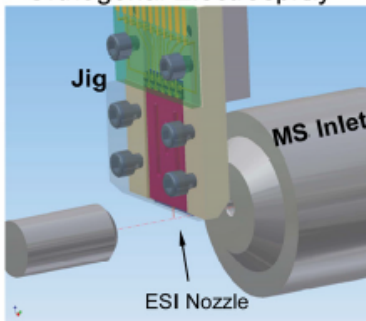


Chip Assembly and Testing Jig

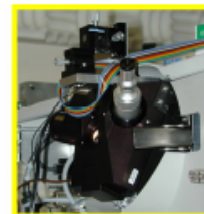


(a)

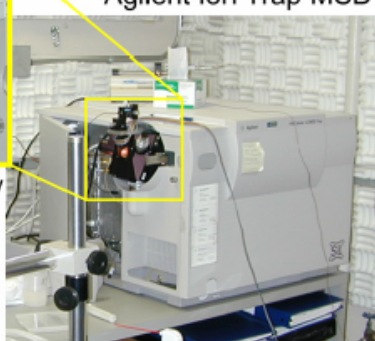
Orthogonal Electrospray



(b)

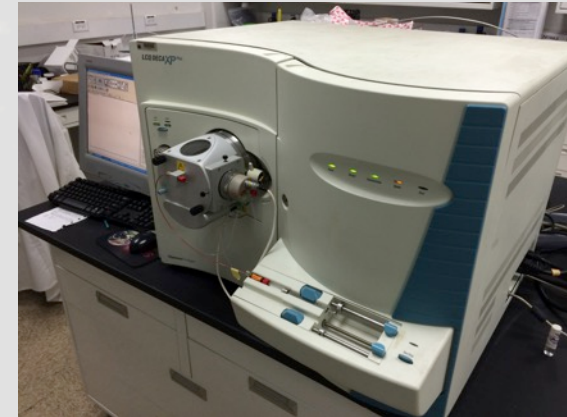
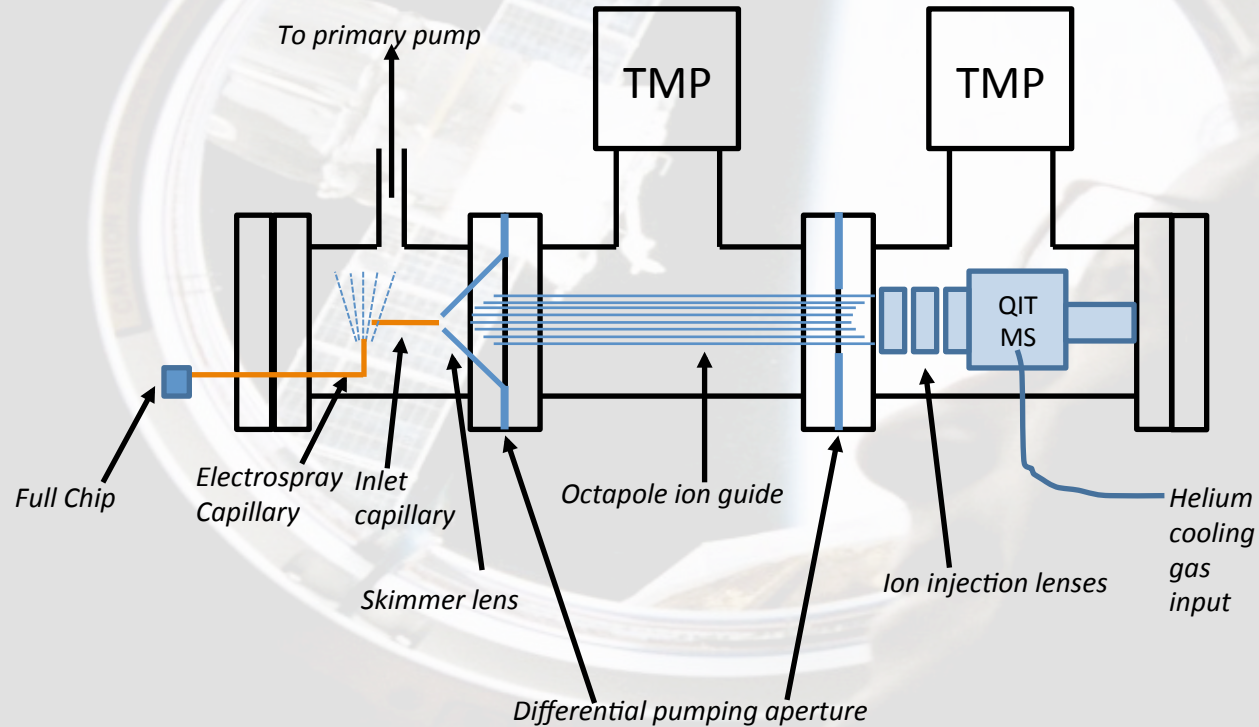


ESI Assembly



Agilent Ion Trap MSD

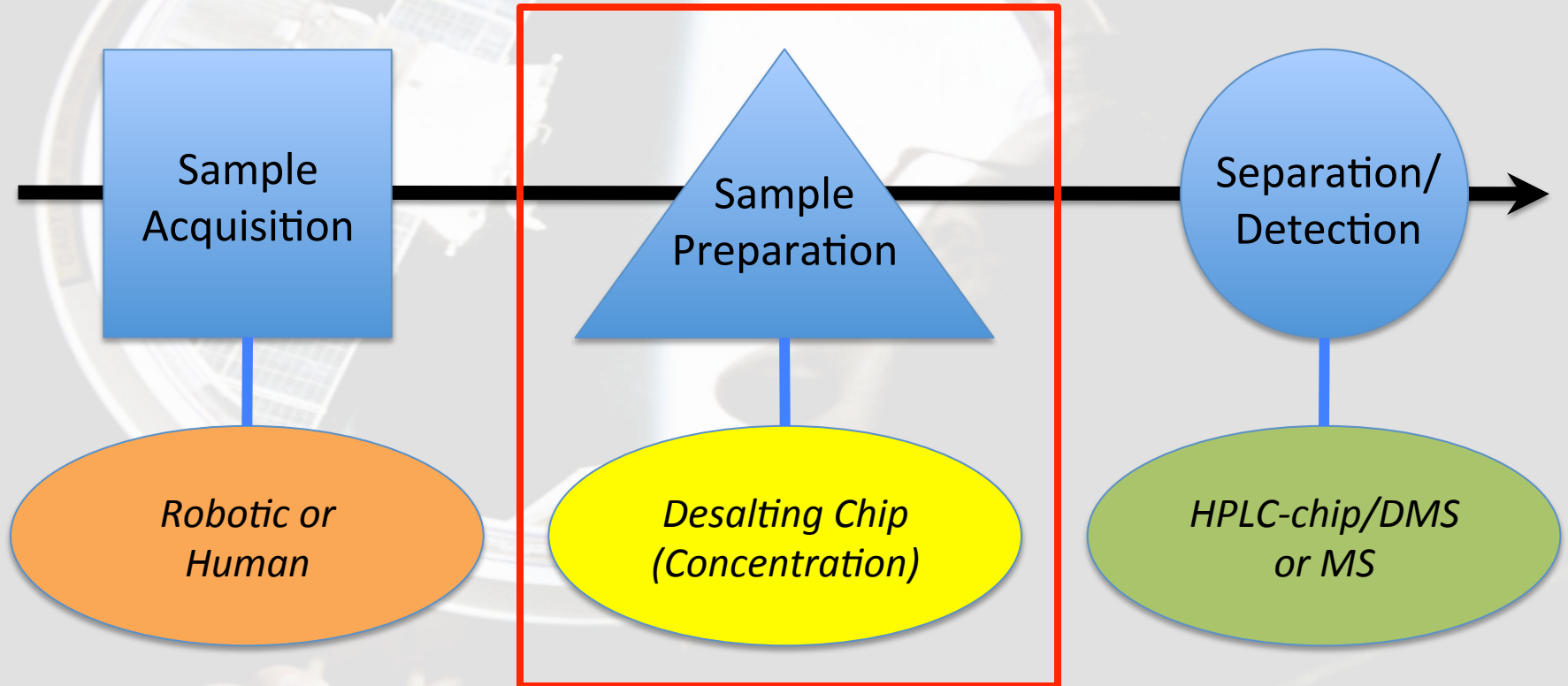
(c)



- Capillary electrospray
- Traditional electrospray in vacuum (Thermo system), minimal pumping needed
- Minimizes vacuum sealing issues on chip side. Safest route



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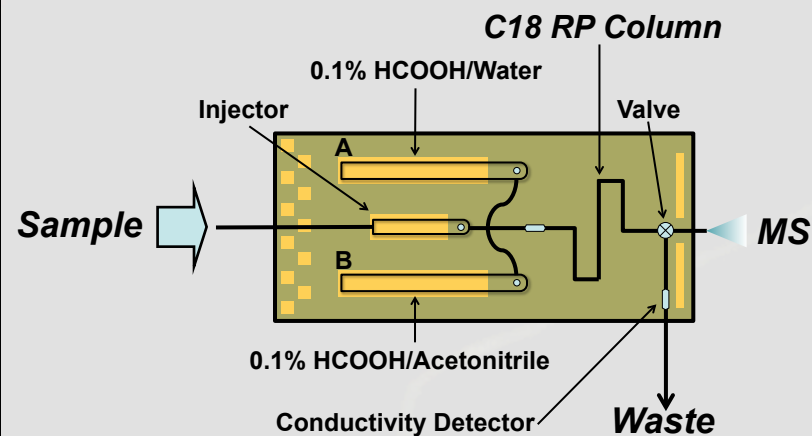
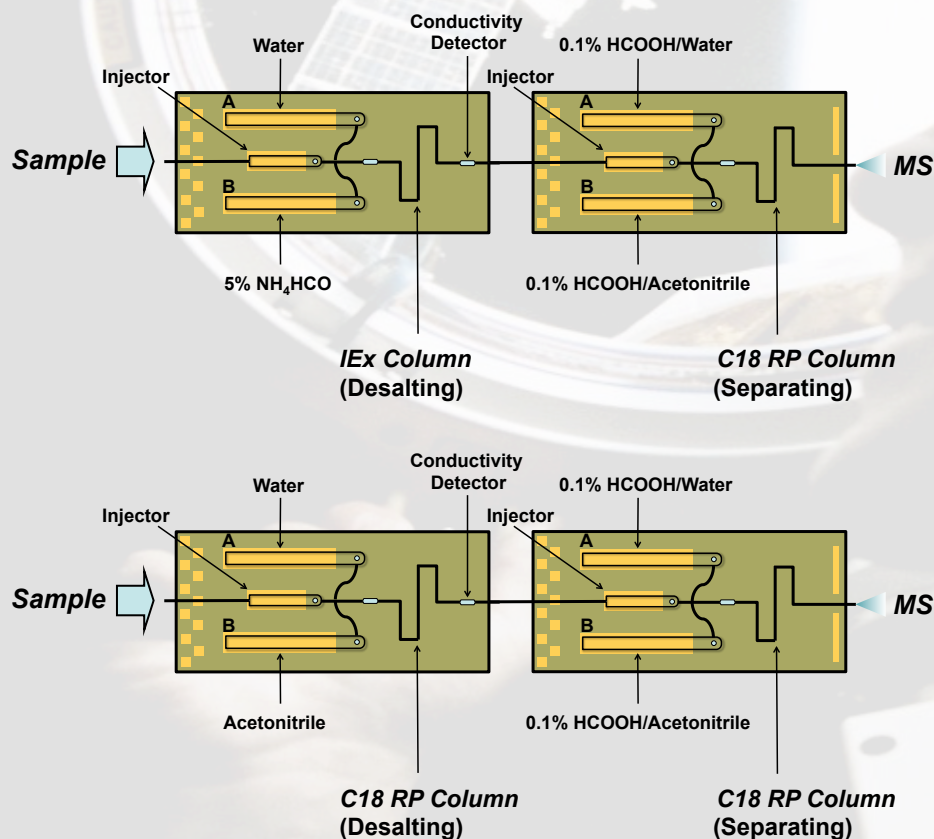


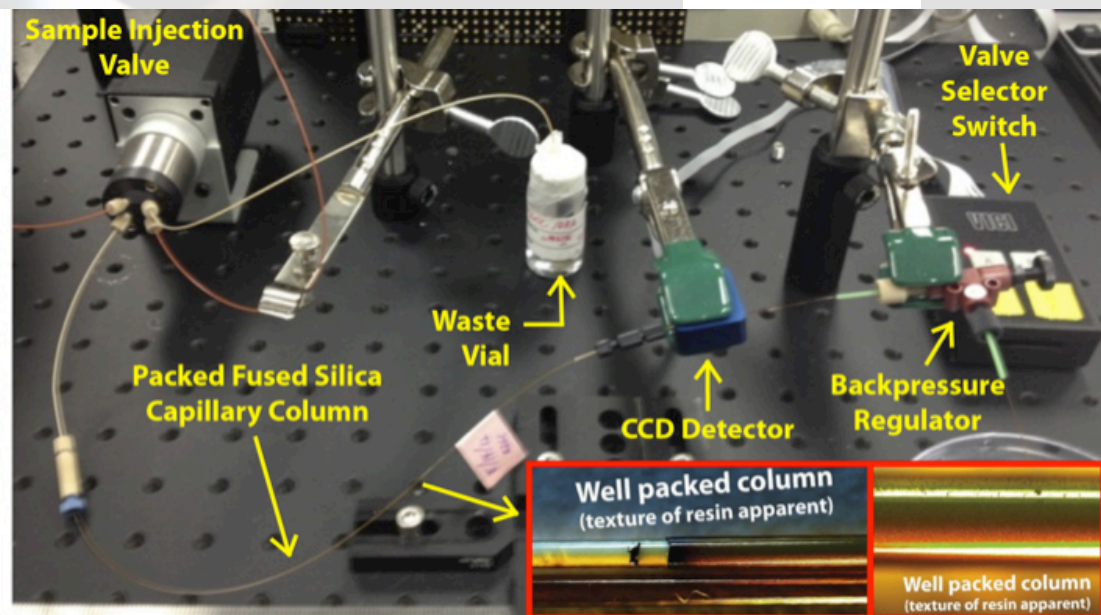
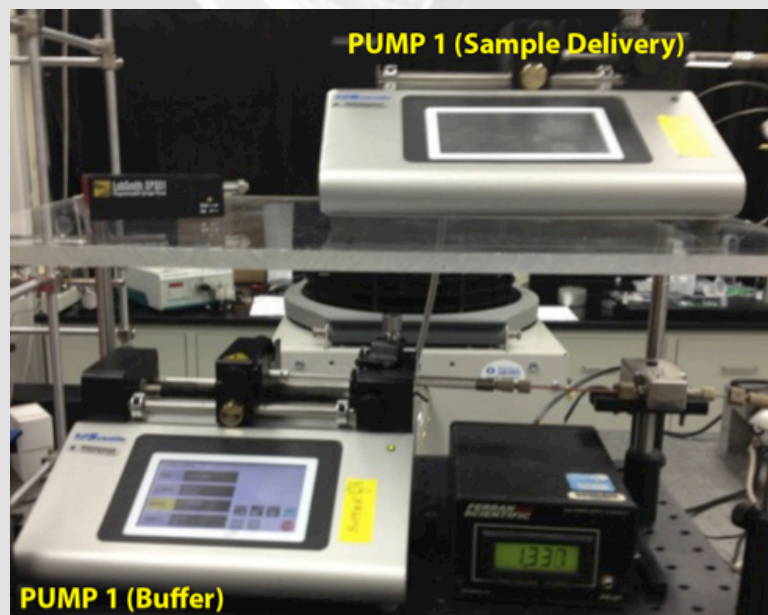
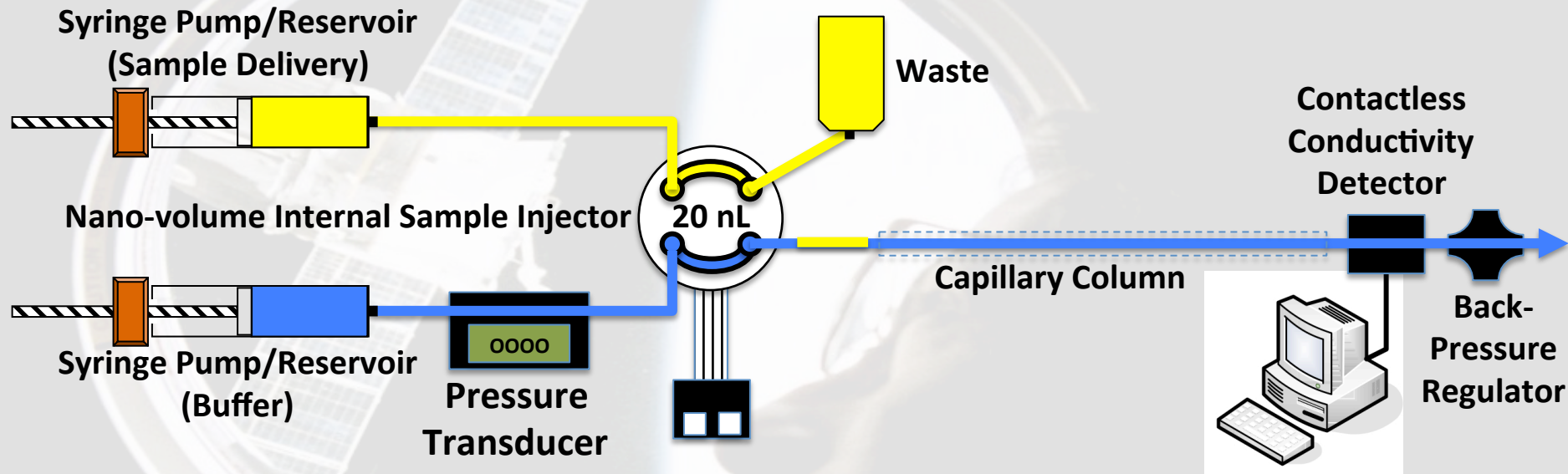
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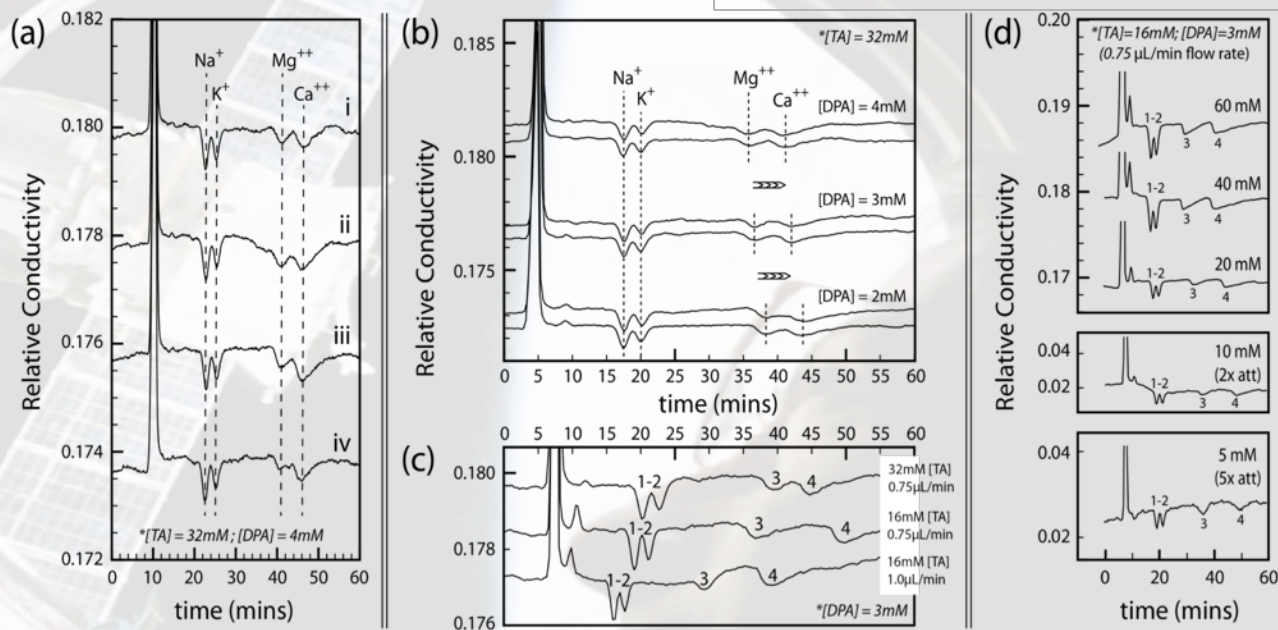




- One step that would be of critical relevance to the HPLC-chip/MS system is desalting; compound mixtures isolated from natural matrices often contain considerable amounts of nonvolatile salts.
- The presence of such salts may interfere with the operation of electrospray ion sources by clogging the skimmer and obscuring or suppressing ionization.
- in complex environments where high levels of salts are present, e.g. Mars (Boynton *et al.*, 2009; Hecht *et al.*, 2009), front-end devices may not be efficient enough (or dedicated enough) to desalt a sample for HPLC/MS.



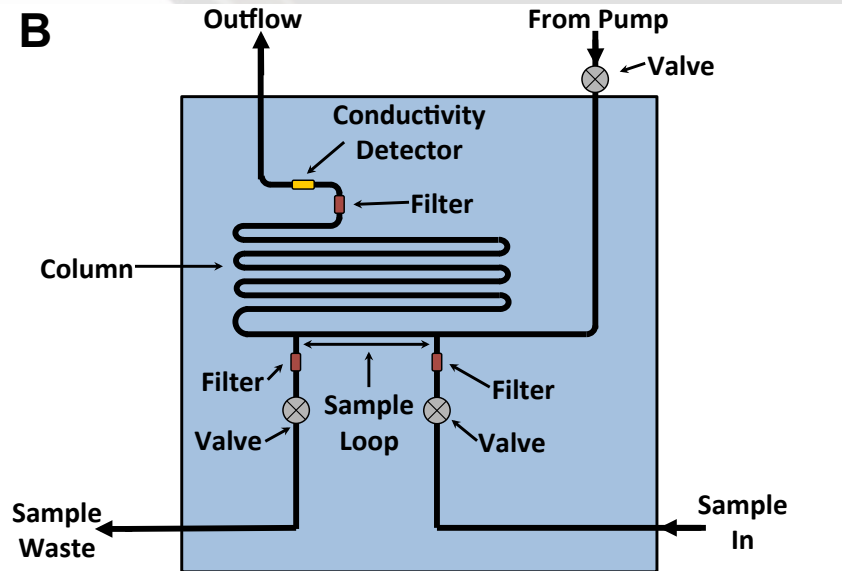
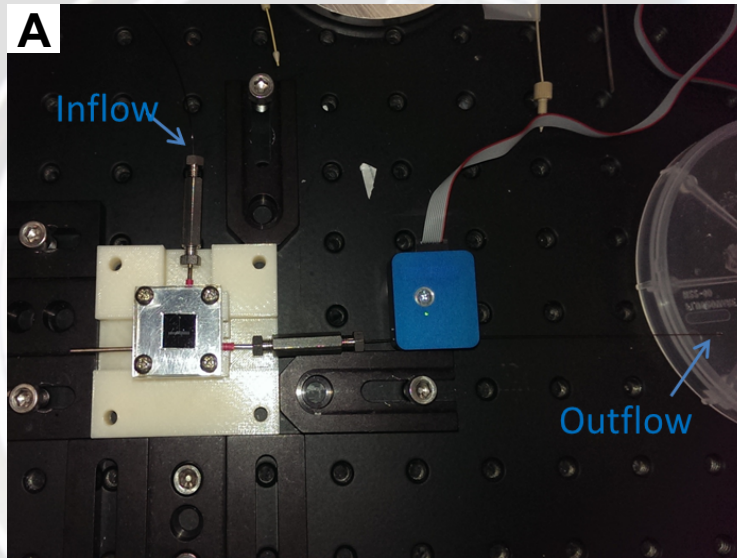




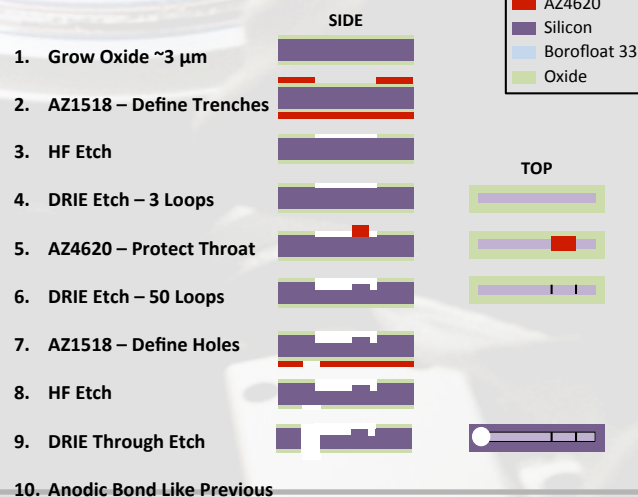
**Test runs conducted to optimize separation conditions.** All experiments used tartaric acid (TA) as the eluting agent and dipicolinic acid (DPA) as a divalent complexing agent; 4-species mixtures were used for these experiments (Na<sup>+</sup>, K<sup>+</sup>, Ca<sup>2+</sup>, Mg<sup>2+</sup>). Capillary columns were packed with 30  $\mu$ m diameter Phenomenex weak cation exchange resin, and pressures were between 0.7-1 MPa. Note that all runs include a system peak before any of the target cations appear – this feature is consistent with the literature.

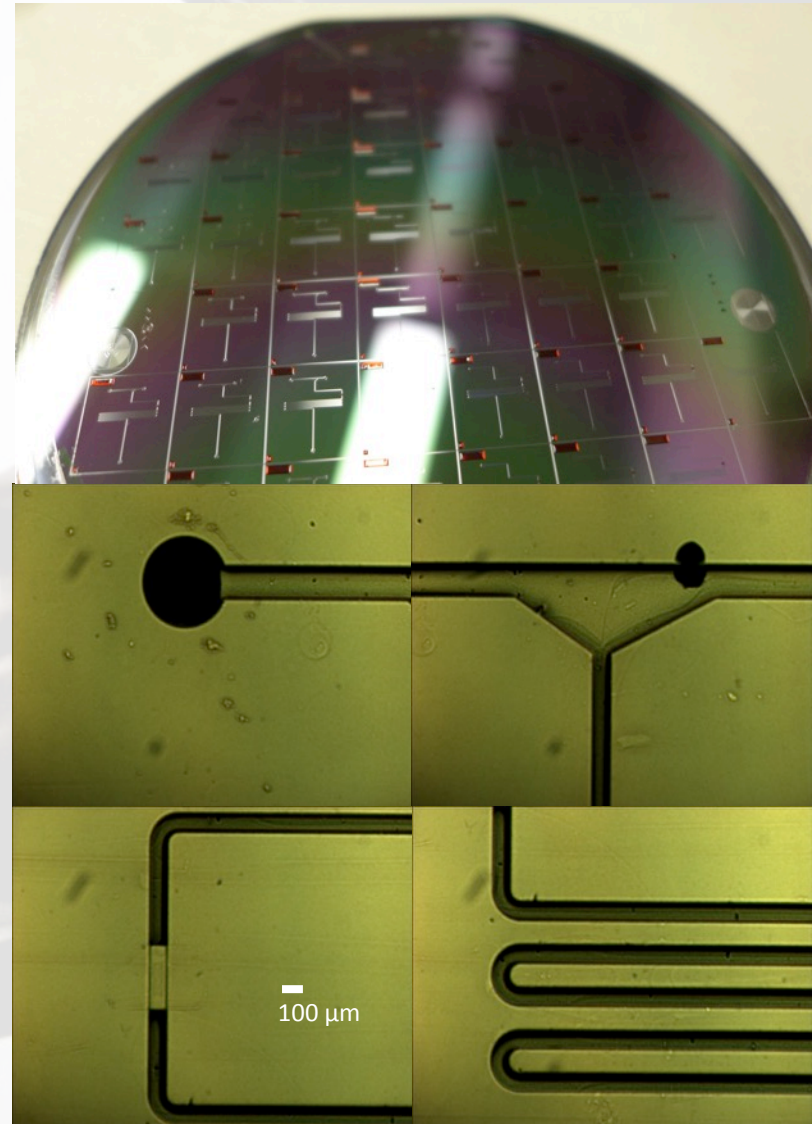
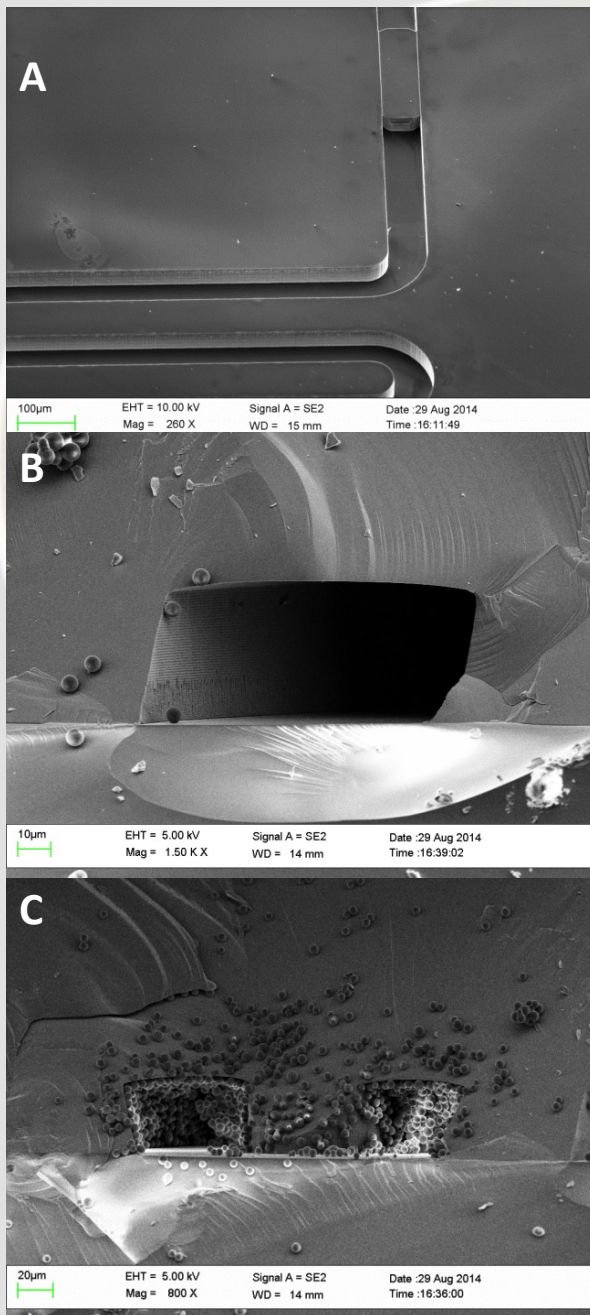
- (a) Reproducibility tests (i – iv): four species resolved at 10 mM concentration (0.5  $\mu$ L/min).
- (b) Effect of [DPA] variability – increased concentrations of DPA reduce divalent retention times (0.5  $\mu$ L/min).
- (c) Effect of [TA] variability – shorter retention times are observed for divalent cations at higher [TA] at flow rates of 0.75  $\mu$ L/min; a similar effect can be achieved using faster flow rates at lower [TA] (in this case 1.0  $\mu$ L/min). While the run at 1.0  $\mu$ L/min is shorter than the 0.75  $\mu$ L/min run by ~10 minutes, the peak resolution benefits are minimal.
- (d) Serial dilution from 5 mM to 60 mM demonstrating detection thresholds in the micromolar range based on the signal-to-noise ratio of the 5 mM run. The sensitivity of the system is not as high as expected.





## Fabrication







## Acknowledgments

- Sabrina Feldman (JPL)
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- Stojan Madzunkov (JPL)
- Jurij Simcic (JPL)
- Rob Hodyss (JPL)
- Aaron Noell (JPL)
- Guyane Kazarians (JPL)
- Andrew Aubrey (JPL)
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- Nicholas Scianmarello (Caltech)

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